



This listing of claims will replace all prior versions of the claims in the application:

**Listing of Claims**

1. (canceled)
2. (canceled)
3. (canceled)
4. (canceled)
5. (canceled)
6. (canceled)
7. (amended) A method for identifying cis-aminoacylating catalytic RNA molecules comprising the steps of:
  - a. providing a tRNA-like molecules;
  - b. providing a ribozyme domain molecules;
  - c. attaching the ribozyme domain molecules to the 5' end of the tRNA-like molecules to obtain a pool of ribozyme-tRNA molecules;
  - d. contacting the ribozyme-tRNA molecules with an amino acid substrate wherein the amino acid substrate is not conjugated to RNA to obtain aminoacylated RNA molecules and non-aminoacylated RNA molecules; and
    - e. ~~partitioning the aminoacylated ribozyme-tRNA molecules from the remainder of the ribozyme-tRNA molecules to obtain cis-aminoacylating catalytic RNA molecules~~  
partitioning aminoacylated RNA molecules from non-aminoacylated RNA molecules wherein the aminoacylated RNA molecules are cis-aminoacylating catalytic RNA molecules aminoacylated with the amino acid substrate at the 3' end of the ribozyme-tRNA molecules.
8. (canceled)
9. (canceled)
10. (canceled)
11. (canceled)
12. (canceled)

13. (amended) The method of claim 7, wherein the tRNA-like molecule consists of SEQ ID NO: 16.
14. (previously presented) The method of claim 7, wherein the ribozyme domain molecule consists of SEQ ID NO: 9.
15. (previously presented) The method of claim 7 wherein the cis-aminoacylating catalytic RNA molecules consist of, from 5' to 3', SEQ IN NO: 9, SEQ. ID NO: 16.
16. (previously presented) The method of claim 7 wherein the amino acid substrate is an N-biotinyl-L-aminoacyl-cyanomethyl-ester.
17. (amended) A method for ~~constructing~~ identifying cis-aminoacylating catalytic RNA molecules comprising the steps of:
- a. providing RNA molecules having a tRNA-like domain and a 5'-leader ribozyme;
  - b. contacting the RNA molecules with an amino acid substrate wherein the amino acid substrate is not conjugated to RNA; and
  - c. partitioning aminoacylated RNA molecules from non-aminoacylated RNA molecules wherein the aminoacylated RNA molecules are cis-aminoacylating catalytic RNA molecules aminoacylated with the amino acid substrate at the 3' end of the ribozyme-tRNA molecules.
18. (amended) The method of claim 17 wherein the tRNA-like domain is SEQ ID NO: 16.
19. (amended) The method of claim 17 wherein the 5'-leader ribozyme is SEQ ID NO: 9.
20. (previously presented) The method of claim 17 wherein the amino acid substrate is an N-biotinyl-L-aminoacyl-cyanomethyl-ester.